

## SEQUENCE LISTING

- <120> METHOD FOR PREPARING 1,3-PROPANEDIOL BY A RECOMBINANT MICRO-ORGANISM IN THE ABSENCE OF COENZYME B12 OR ONE OF ITS PRECURSORS
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- <140> 10/043,639
- <141> 2002-01-09
- <150> PCT/FR00/01981
- <151> 2000-07-07
- <150> FR 99/08939
- <151> 1999-07-09
- <160> 10
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<b>-1</b> -	<b>a</b> 1	ml	77-	<b>M</b>	m	17	2	0	<i>α</i> 1	C	17- 7	T	T	17- 1	тіс
шe	Glu	Thr	Ala	мет 165	Tyr	vaı	Asn	ser	170	ser	vaı	гÀг	гуѕ	Val 175	11e
				105					1,0					175	
Pro	Tyr	Ile	Asp	Leu	Ala	Met	Ile	Asp	Ile	Lys	Ser	Met	Asn	Asp	Glu
			180					185					190		
						_	_								
Ile	His	_	Lys	Phe	Thr	Gly		Ser	Asn	Glu	Ile		Leu	Gln	Asn
		195					200					205			
Ile	Lys	Leu	Ser	Asp	Glu	Leu	Ala	Lys	Glu	Ile	Ile	Ile	Arg	Ile	Pro
	210			•		215		•			220		_		
	Ile	Glu	Gly	Phe		Ala	Asp	Leu	Gln		Ile	Gly	Ala	Ile	
225					230					235					240
Gln	Phe	Ser	Lars	Ser	Leu	Thr	Asn	Len	Lve	Ara	Tle	Asn	Leu	Leu	Pro
U 111	1110		- Y - 2	$\mathcal{L}_{L}$	L C U	T 11T	11011	عات لا	-13	٠ ٠ - ٩		٠.5٢	_ u		

Tyr His Asn Tyr Gly Glu Asn Lys Tyr Gln Ala Ile Gly Arg Glu Tyr
260 265 270

Ser Leu Lys Glu Leu Lys Ser Pro Ser Lys Asp Lys Met Glu Arg Leu 275 280 285

Lys Ala Leu Val Glu Ile Met Gly Ile Pro Cys Thr Ile Gly Ala Glu 290 295 300

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<211> 385

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<213> Clostridium butyricum

<400> 8

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20 25 30

Lys Ala Leu Ile Val Thr Asp Lys Phe Leu Lys Asp Met Glu Gly Gly 35 40 45

Ala Val Glu Leu Thr Val Lys Tyr Leu Lys Glu Ala Gly Leu Asp Val
50 55 60

Val Tyr Tyr Asp Gly Val Glu Pro Asn Pro Lys Asp Val Asn Val Ile 65 70 75 80

Glu Gly Leu Lys Ile Phe Lys Glu Glu Asn Cys Asp Met Ile Val Thr
85 90 95

Val Gly Gly Ser Ser His Asp Cys Gly Lys Gly Ile Gly Ile Ala 100 105 110

Ala Thr His Glu Gly Asp Leu Tyr Asp Tyr Ala Gly Ile Glu Thr Leu 115 120 125

Val Asn Pro Leu Pro Pro Ile Val Ala Val Asn Thr Thr Ala Gly Thr 130 135 140

Ala Ser Glu Leu Thr Arg His Cys Val Leu Thr Asn Thr Lys Lys

145 150 155 160

Ile Lys Phe Val Ile Val Ser Trp Arg Asn Leu Pro Leu Val Ser Ile
165 170 175

Asn Asp Pro Met Leu Met Val Lys Lys Pro Ala Gly Leu Thr Ala Ala 180 185 190

Thr Gly Met Asp Ala Leu Thr His Ala Ile Glu Ala Tyr Val Ser Lys 195 200 205

Asp Ala Asn Pro Val Thr Asp Ala Ser Ala Ile Gln Ala Ile Lys Leu 210 215 220

Ile Ser Gln Asn Leu Arg Gln Ala Val Ala Leu Gly Glu Asn Leu Glu 225 230 235 240

Ala Arg Glu Asn Met Ala Tyr Ala Ser Leu Leu Ala Gly Met Ala Phe 245 250 255

Asn Asn Ala Asn Leu Gly Tyr Val His Ala Met Ala His Gln Leu Gly 260 265 270

Gly Leu Tyr Asp Met Ala His Gly Val Ala Asn Ala Met Leu Leu Pro 275 280 285

His Val Glu Arg Tyr Asn Met Leu Ser Asn Pro Lys Lys Phe Ala Asp 290 295 300

Ile Ala Glu Phe Met Gly Glu Asn Ile Ser Gly Leu Ser Val Met Glu 305 310 315 320

Ala Ala Glu Lys Ala Ile Asn Ala Met Phe Arg Leu Ser Glu Asp Val 325 330 335

Gly Ile Pro Lys Ser Leu Lys Glu Met Gly Val Lys Gln Glu Asp Phe 340 345 350

Glu His Met Ala Glu Leu Ala Leu Leu Asp Gly Asn Ala Phe Ser Asn 355 360 365

Pro Arg Lys Gly Asn Ala Lys Asp Ile Ile Asn Ile Phe Lys Ala Ala 370 375 380

Tyr

385

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      Primer
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OIPE

# RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/043,639A

DATE: 04/24/20:13

TIME: 16:21:33

Input Set : D:\Chep004.app

Output Set: N:\CRF4\04242003\J043639A.raw

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3 :110 · APPLICANT: SARCABAL, PATRICIA
              CEOUX, CHEISTIAN
              SOUCAILLE, PHILIPPE
       <120 - TITLE OF INVENTION: METHOD FOR PREPARING 1,3-PROPANEDIOL BY A RECOMBINANT</p>
              MICRO-ORGANISM IN THE ABSENCE OF COENZYME B12 OR ONE OF
              ITS PRECUESORS
     11 - 130 - FILE REFERENCE: CHEP: CO4US
     1: -1140 - CURRENT APPLICATION NUMBER: 10/043,639A
C--> 14 <141> CURRENT FILING DATE: 2003-04-12
     16 H150 - PRIOF APPLICATION NUMBER: PCT/FR00/01981
     17 - 151 - PRIOR FILING DATE: 2000-07-07
     19 -150 - PRIOR APPLICATION NUMBER: FR 99/03939
     20 K151 - PRIOF FILING DATE: 1999-07-09
     II + 160 - NUMBER OF SEQ ID NOS: 10
                                                            ENTERED
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     De - 210 - SEQ ID NO: 1
     .7 H.111 LENGTE: 2364
     The HOlds TYPE: DNA
     19 -0115 - OFGANISM: Clostridium butyricum
     31 -4400 > SEQUENCE: 1
     30 utgahaagta aaggalittag taoocaaaca gaaagaataa atattttaaa ggotoaaata 🍪
     Po tisaatgota aascatgigi igaatoagaa agagcaatai taataacaga atcatttaaa 170
     34 masamagaag gomagobago sattttaaga agagoattgg cattgaaaca catabttgaa 186
     35 uataboodta taadaattag agabdaagaa obtatagtgg gaagbbbaad baaagaadda 240
     36 Sygtettess aagtauttee teagtittiet aataagtegt tacaagatga attegataga 300
     27 ttaabtaaga gaustggaga tgoattosaa atttoagaag saagtaaaga aaaattaasa 500
     36 yatgtottty agtatrygaa tygaaagaca abaagtgagt tagbaactto atatatgaca 410
     39 magguaadaa gamaggdagt aaattigtigaa gtatttadtig taggaaacta otattataat 480
     40 ggognaggad atqtatotgt agantlatgga aaagtattaa gggbtggatt taatgggatt 540\,
     41 atawatgagg otwaggwaca attagawwaa wacaggagta tagatootga tittatawag 600
     4% aaagkaaaat tootakatag tittattato toatgogaag otgoaataac atatgtaaat 660
     45 agatatgota aawaggotaa agagattgoa yataatabaa gtgatgoaaw aagawaagst 710
     44 gaartaaaty aawtaqbaaa aattigtica saagtiibag gagagggage taaatciitte 780
     45 tatqaagcat gicaafitatt tiggittatt batgcaataa taaatataga atolaatgga 840
     40 patietatit etopagetag attidateaa tabatgiate eatattatga aaatgataaa 900
     47 aatataacag atkaagitigo toaaqaatta atagatigta totggattaa attaaatgat 960
     48 attaataaag taagagatga gatttoaact aaacattttg gtggttaccc aatgtatcaa 1020
     49 aaattaattig ttigiggijtea aaatteaqaa ggaaaagatig caactaataa agtateatat 1080
     50 atggcattag aagsagetgt eeatqtaaag tigeetcags catetitigte agtaagaata 1140
     51 tygaataaqa otobagatga attittigott agagoagoag aattaaotau agaaqqytta 1700
     52 ggaetteotg ettathataa tgatgaagtt attahteead cattagttte tadaggtott 1770
     53 apattagaag atgcaayaga otabygaata attggatyty ttyaabcaba aaagcbagga 1520
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54 aaaacagaag gatggbatga tibarhattu titaatotid caadaatagt adagttaact 1390

PATENT APPLICATION: US/10/043,639A TIME: 16:21:33

DATE: 04/24/2003

Input Set : D:\Chep004.app

Dutput Set: N:\CRF4\04242003\J043639A.raw

```
E5 ataaattotg gatttgataa aaataaabag attggabota aaabtbaaaa ttttgaagaa 1443
-6 atgaaatoot itgatgaatt batgaaagot tataaagoto aaatggagta tittigtaaaa 1800
z^{lpha} catatytyst ytystyataa tiysatayat attysasaty sayaaayays tissattasit 15\,6\%
88 ttottgtbat baatggttga taattgtato ggaaaaggaa agagoottba agatggtggt 1623
69 goagaatata Acticagtgg accacaaggt gitggagtag ctaatatigg agattcatta 1000
60 gttgdagtta aasaasttgt gtttgatgas astaagatta oboottdaga attaaayasa 1.34\%
61 adattaaata atgattitaa aaattoagaa gaaatadaag dottadtaaa aaatgotoot 1863
e. Aagtteggaa aegatattga egaagetgae aatteageta gagagggege attagtatae 1866
n'n typaqaqaag teaataaata tacaaatooa aqqqqaqqaa attibbaabo aqqatbatat 1920
64 opatottoaa ttaatgtata tititggaago ttaacaggitg otactocaga tggaaggaaa 1980
65 thoographic cattagodga tyggggtttot obatbaagag gotgtgatgt atotggabot 7.040
66 actgoagott gtaactcagt tagtaaatta gatcatttta tagcttcaaa tggaacttta 2197
67 titaatoaaa aattooatoo gooagoatta aaaggtgata atggattaat gaatttatoa 2160
66 toattaataa qaaqttatti tgatcaaaay ggatttoatg ticaatttaa tgtaatagat 2.7.0
69 waaaaaatat tacttgcago acaaaaaaat ootgaaaaat atcaagattt aattgttaga 2280
70 gttgcaggat ataqtgcaca gttcatttct ttagataaat ctattcaaaa tqatattatt 2340
                                                                       .15 € 4
VI gdaagaactg aacatgttat gtaa
74 <210 + SEQ ID NO: 2
75 < 211 - LENGTH: 915
76 + 212 > TYPE: DNA
77 - 213 - ORGANISM: Clostridium butyricum
79 + 400 > SEQUENCE: 2
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61 cotqqaataa qaactatagt attttttaaq qqatqttcaa tqtcqtqctt atqqtqcact 110
80 aatodagaat oobaagatat taaadottoaa gtaatgttta ataaaaattt atgtadaaaa 180
65 tqtgqaaqat qtaaatctca atgtaaaaqt gcaggtattg atatgaattc agaatatagg 240
64 atadataasa qossatqtad agaqtqtada asatqtqttq ataattqctt asqcqqqqda 300
85 ottgitatig aaggaaggaa tiacagtgit gaagacgita taaaggaati gaaaaaagat 300
be agtighticast atagaagatic aaacggtigga attacactat ctiggaggiga agtattactt 410
67 daacdagatt tigragigga gottitaaaa gagigtaaat dataiggoig goadacigoo 480
88 attgaaabag caatgtatgt taatagtgaa totgtaaaaa aagtaattoo atatatagat 540
FM ctggctatja ttgatataaa aagtatgaat gatgaaatoo ataggaaatt tacaggagtg 606
🗝 aytaacgasa taatattaca aaacattaaa ttaagtgatg aattagctaa agaaataata 🕬
91 atbaqaatto otgtaataga aggatttaat goagatttab aaagtatagg agbaatagot 700
91 daattittosa aatoattaad aaatottaaa agaatagato tiottodata odataattat 780
43 qqaqaaaata agtatcaago aattggaaga gagtattott tgaaagaact aaaatcacct 840
🥯 aqtawagaca aaatggaaag attaaaagct ttagttgaaa tcatgggaat accgtgcaca 💯
95 attggagetg agtaa
                                                                       915
95 - 110 - SEQ ID NO: 3
93 + 311 + LENGTH: 28
100 -021. > TYPE: DNA
101 - 2130 OFGANISM: Clostridium butyricum
1:3 :40: - SEQUENCE: 3
                                                                        28
1.4 tagamaaaac awacaaaaat gttattat
147 K210 + SEQ ID NO: 4
108 -: 211: LENGTH: 1158
109 - 212: TYPE: DNA
110 - 213: OFGANISM: Clostridium butyricum
112 -400> SEQUENCE: 4
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PATENT APPLICATION: US/10/043,639A TIME: 16:21:33

DATE: 04/24/2003

Input Set : D:\Chep004.app

Output Set: N:\CRF4\04242003\J043639A.raw

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113 atgagaatgt atgattattt agtaccaagt gtaascttta tgggagcaaa ttcagtatca 60
114 graduagety asagatycas astattaggt ggassassag cattgatagt tacagatasg 110
11% tittotaaaag ataiggaagg tiggagoigti gaattaabag tiaaatatti aaaagaagbi 1---
116 ygattagatg tigtatatta tgaoggagit gaaccaaatc caaaagatgi taatgitata 340
11/ qaaqqattaa aaatatttaa aqaaqaaaat tgtqacatqa tagtaactqt aqqtqqaqqa 300
11% Agitingbatg attgoggtaa gggaatagga attgotgbaa babatgaagg agatotttat 160%
119 quitatgrag gaatagaaac actigicaat coattgocac caatagtage tigtaaatact 400
120 actiquaggaa otgotagtiga attaactogt battiqtigtat tigabtaatab aaaaaaagaaa 4---
121 araawattig tiatagitag oliggagaaat tigoototag talotalaaa igalocaatg (40
1.00 ottatggtos sassacotgo aggattasca gosgotacag gastggatgo tittascacat 600
10% gwaatagaag catatgtato aaaagatgoa satocagtaa cagatgotto agcastacaa 660
124 getattaaat taattttoaca asatttaaga caagetgtag otttaggaga asatottgas 7.0
155 gbasyagaaa atatggotta tigbatbatta otagbaggaa tigibatttaa taatgotaat 780
126 thaqqatatg tacatgcaat ggotcatcaa ttagggggac tgtatgatat ggcacatggt 840
1.17 quidetaatg caatgetatt accadatgit gaacgitata atatgetate aaateetaag 900
11.3 aagtittgdag atatagdaga attitatggga gaaaatatat otggactitto tqtaatggaa 960
110 qoaqeaqaqa aaqeeataaa tqeaatqtto aqqetttoaq aqqatqttqq aattooqaaa 10.0
130 aqtotaaaqq aqatqqqaqt qaaacaaqaa qattttqaqc atatqqcaqa actaqctott 1080
171 ttagatggaa atgootttag caatocaaga aaaggaaatg caaaagatat tataaatatt 1140
132 Uttaaggotg ottattaa
1:5 K210 - SEQ ID NO: 5
136 4011 LENGTH: 4963
137 HU12 - TYPE: DNA
135 - 0.015 - ORGANISM: Clostridium butyricum
140 - 400 - SEQUENCE: 5
141 qaataaaagt tatotataaa tgataaaagt battattaga taacttitta tiittaaaaata 🕬
142 abtactaata aasagttosa agastattac agtagacatt tgasagasatg castgatasa 1.0
145 caattgtatt agstittaact itagataaaa caaacaaaaa tgittattatt agccaagaaa 180
144 atactgttac aaaagaaaag agaaaaacat agcaaaagag taccaatatt aagcaataaa 040
145 mittottaaa atattatosa tassaigata agattagata aabbaagtaa gastgigatt 500
140 dyaggagtaa aaatgataag taaaggatti agtabobaaa bagaaagaat aaatattita 😿 🖰
147 Haggetbaaa taltaaatge taaaccatgt gttgaatcag aaagagcaat attaataaca 4.0
144 quatcattta aacaaacaga aggocagoca gcaattttaa qaaqagcatt ggcattqaaa 440
149 cacatactig aawatatooc tataacaatt agagatcaag aacttatagt gggaagttta 540
150 actaaagaad cagatgotto acaagtattt ootgagtttt otaataagtg gttacaagat \psi\psi\psi
151 gaattggata gattaaataa gagaactgga gatgcattoo aaatttaaga agaaagtaaa 660
152 gaaaaattaa aayatgtott tgagtattgg aatggaaaga caacaagtga gttagcaact 520
15% thatatatga baqagyaaab aaaagatgoa gtaaattgtg aagtatttab tgtaggaaab 2%.
154 tactattata atigograph acathtatet grapattath gaaaagtatt aagggtigga 840
15^{\circ} thiaatggga trataaatga ggotaaggaa caattagaaa aaaacaggag tatagatoot 96^{\circ}
15) gattitataa agaaagaaaa attootaaat agtgitatta totoatgoga agotgoaata 9x0
11\% anataigtaa atagatatyo taaaaaaggot aaagagatty cagataatac aaaagatyoa 101.0
18% awaagaaaag otgaattawa tgaaatagca aaaatttgtt caaaagatac aggagaggga 10%0
189 gitagatott totatgaago atgtoaatta tittggttta tacatgoaat aataaatata 1940
100 quateraatg gasattotat tiotocagot agattigato aatocagtaa tocatattat 1200
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162 amattabatg atattaataa agtaagagat magatttoaa ptamacattt tgytgyttac 1320
163 chiatgiato aaaaattaat tyttyyyyyt saasattoag aagyaaaaga työäästaat 1380\,
164 awagtatoat atatggottt agaagoagot gtowatgtaa agttgootea gooatetttg 1440
```

PATENT APPLICATION: US/10/043,639A TIME: 16:21:33

DATE: 04/24/2003

Input Set : D:\Chep004.app

Output Set: N:\CRF4\04242003\J043639A.raw

165 teagtaagaa tatggaataa gaeteeagat gattttgage ttagageage aggattaaet 1500 166 agagaayggt taggacttoc tycttattat satgatgaag ttattattoc agcattagit 156% $16^\circ$  totagaggto thacattaga atatagbaga gabtaoggaa taattggatg tgttgaacba 16.00168 baaaagobag gaaaaabaga aggatgybat gattatgbat totttaatot tyaaagaata 168: 16 my gragagotaa obabaaaboo tggaboogab aaaaaaagaab agaboggado baaaabooaa 1 % my 17) aattittgaag aaaggaaato ottitgatgaa tioatgaaay ottataaago toaaatiggag 1800 1% tatitiqiaa aabataiqig biqiqotdat aaatqataaq atatiqiada tqoaqaaaqa  $1/m_{\odot}$  $^{\circ}$ .) gotocattas otttottigto accapatigti gataattigta toygaaaagg aaagagcaat 1.6.91% baagstytag gtgbagaata taabttbagt ggalbabaag gtgttygagt agetaatatt  $1 \ll 1$ 1.4 ggagattbat tagttgbagt taaaaaaaatt gtgtt:gatg aaaataagat tabtbbtbba 2.0491°5 gaattaaaga aaabattaaa taatgattit aaaaattoag aagaaataca agoottacta 🗈 176 aaaaatgoto obaagtotgg aaatgatatt gatgaagtog ataatotago bagagagggt 11601%% goattagtat actgtagaga agttaataaa tatacaaato caaggggagg aaattttcaa 1.7.017% obaggattat atobatotto aattaatgta tattttggaa gottaabagg tgotabtoba  $\mathbb{R}^{2,0}$  $1^\circ$  4 gatggaagga aatooggada abbattagot gatggggttt btobatcaag aggotgtgat .54915% gtatotygas obactycago tigitaactoa gibagtaaat bagatoatti batagottoa 34%160 aatggaactt tatttaatca aaaattocat oogtoagoat taaaaggtga taatggatta .400lfo atgaatttat patpattaat aagaagttat titgatpaaa agggatitoa tyttpaattt  $\mathcal{AL} \Phi$ 18) aatguastag ataaaaaaat attaottgoa goabasaaaa atootgaaaa atatoaagat  $\mathbb{M}^{4,6}$ 164 theattqtta gagttgcagg atatagtgca cagttcattt otttagataa atctattcaa .040 188 aatgatatta tigoaagaab tgaabatgit atgiaaagab agbiittaaa ggggataaaa 120018% ghaangagna aggagahaaa aggoghtiba bhisaabahab aaaaabbibo ghbabahgab 17%%167 gggeetggaa taagaaetat agtattittt aagggatgtt caatgtogtg ottatggtgolik. 154 agtaatobag aatobbaaga tattaaaoot baagtaatgt ttaataaaaa tttatgtaba 2000 16 + adatgtgggaa gatgtaaato toaatgtaaa agtgoaggta ttgatatgaa ttoagaatat 104018% aggatagata asagcasaty tabagaytyt acasasatyty ttystaatty ottaagogyy 800%191 gbabbgtba bbgaaggaag gaabbabagb gbbgaagabg bbabaaagga abbgaaaaaa 5000 1% gatagtiztic satisfaced atcassogic ggasttaced tatotygazg ggasztatta 51%193 obticallocal attitiguage ggagottita alaqaytyta alatcatatgy otygologica 318%194 godattgaaa bagbaatgta tgttaatagt gaatotgtaa aaaaagtaat tobatatata 3340 19/ gatotgotta tgattgatat aaaaagtatg aatgatgaaa tooataggaa atttacagga x+1 / 196 gtgagtaacg aaataataatt acaaaaacatt aaattaagtg atgaattago taaagaaata 3566 ataatbagaa tibotgiaat agaaggatti aatgbagatt tabaaagtat aggagbaata 🖽 🕡 198 gotoaattit baaaatoatt aabaaatott aaaagaatag atottottoo atabbataat 5485 13 ) batggagaaa ataagtatoa agoaattgga agagagtatt otttgaaaga actaaaaatca  $\pm 64\%$ 200 obtaigtawag wasawatiga wagattawa gotttagttig waatbatiggi wataabigtiga 500%261 abaattiggag otgagtaata gtagotttab abbagatatt ttaaaaabaa ttttaaaatta  $\phi \phi \phi$ 200 awaggagaag attgoatatg agaatgtatg attatttagt abbaagtgta aabtttatgg  $\pm700$ 200 gagoaaatto agtatoagta gtaggtgaaa gatgoaaaat attaggtgga aaaaaaagcat  $\pm 780$ 204 tgatagttas agataagtti otaaaaagata tggaaggtig agotigtigaa itaacagtta 3840 205 aatattiaaa agaayotgga ttagatgttig tatattatja bagagttaaa ocaaatobaa 3900 200 aagatgitaa tgiratagaa ggattaaaaa tatttaaaya ayaaaattgi gacatgatag 3960 207 taactgtagg tggaggmagt togcatyatt gryytaagyg aataggaatt getgoaacab 4000 208 atgaadgaga totttatgat tatgbaggaa tagaaadabt tqtcaatoca ttqccaccaa  $40^{4}0$ 209 tagtagotut saanabtabt goaggaacty otagtgaant aactogtoat tyngtattya 4140 210 otaatacaaa saagaasaha asattigita tagittagoig gagasahhig ocholaghat 4200 211 otataaatga toosatjott atggtesass sacetquagg attaadajis gmanaggas 4200 212 tygatootti aanapatoon atagaaboat atgiafinkaa adatgisaakti cicagtaabay 4320 III atgottoano asticaadon attaaattaa titoacaaaa titaaadacia gotgiinotti 430

DATE: 04/24/2003 PATENT APPLICATION: US/10/043,639A TIME: 16:21:33

Input Set : D:\Chep004.app

Dutput Set: N:\CRF4\04242003\J043639A.raw

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010 atgatatggo adatggtgtt gotaatgdaa tgotattado adatgttgaa ogttataata 4500
17 tgotatcaaa tootaagaag titigoagata tagoagaatt tatgggagaa aatatatotg 4000
118 gaotttotqt aatggaagea qoaqaqsaag obataaatgo aatgttoagg otttoagagg 4880
21→ atqttqqaat tooqaaaaqt otaaaqqaqa tqqqaqtqaa acaaqaaqat tttqaqcata 4740
120 tygoagaact agototttta gatygaasty ootttagoaa tooaagaaaa yyaaatyoaa 4800
2.1 aagatattat aastattttt aaggetgett attaattaat aetatttaaa ggatteaaag 4860
Lii taaaagataa aadatatata tattagatti aagattttat tataggotaa caacaaagaa 4920
123 haagttaagt attaaastta gottgttott tgttgtttat ttt
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327 -211> LENGTH: 783
22% - 112> TYPE: PRT
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131 <400> SEQUENCE: 6
THE Met Ile Ser Lys Gly Phe Ser Thr Thr Glu Arg Ile Asr. Ile Leu Lys
    1
                                         10
225 Ala Gln Ile Lou Ash Ala Lys Pro Cys Val Glu Ser Glu Arg Ala Ile
                                     25
                ΞÚ
                                                         31)
23% Leu Ile Thr Glu Ser Phe Lys Gln Thr Gly Gln Pro Ala Ile Leu Arg
                                 40
                                                      4 =
241 Arg Ala Leu Ala Leu Lys His Ile Leu Glu Asn Ile Pro Ile Thr Ile
242
        50
                             5.5
                                                  ნ∷
144 Arg Asp Gln Glu Leu Ile Val Gly Ser Leu Thr Lys Glu Pro Arg Ser
245 65
                         70
                                             75
147 Ser Gln Val Phe Pro Glu Phe Ser Ash Lys Trp Leu Gln Asp Glu Leu
                                         90
0.4\,\mathrm{H}
                     85
250 Asp Arg Leu Ash Lys Arg Thr Gly Asp Ala Phe Gln Ile Ser Glu Glu
251
                                    105
253 Ser Lys Glu Lys Leu Lys Asp Val Phe Glu Tyr Trp Ash Gly Lys Thr
                                120
            115
Not The Ser Glu Leu Ala The Ser Tyr Met The Glu Glu The Arg Glu Ala
. 5.7
                            135
                                                 140
159 Val Ash Cys Glu Val Phe Thr Val Gly Ash Tyr Tyr Tyr Ash Gly Val
160 145
                        150
                                            155
101 Gly His Val Ser Val Asp Tyr Lys Val Leu Arg Val Gly Phe Asr. Gly
                                        170
                    165
If: The The Ash Glu Ala Lys Glu Gln Leu Glu Lys Ash Arg Ser Asp Pro
266
                180
                                    185
                                                         190
2005 Asp Phe Ile Lys Lys Glu Lys Phe Leu Asn Ser Val Ile Ile Ser Cys
269
           195
                                200
                                                     205
171 Glu Ala Ala Ile Thr Tyr Val Asn Arg Tyr Ala Lys Lys Ala Lys Glu
                            215
                                                 220
        210
274 Ile Ala Asp Asn Thr Ser Asp Ala Lys Arg Lys Ala Glu Leu Asn Glu
27! 2:5
                        230
                                             235
2"" Ile Ala Lys Ile Cys Ser Lys Val Ser Gly Glu Gly Ala Lys Ser Phe
                                        250
                    245
280 Tyr Glu Ala Cys Gln Leu Fhe Trp Phe Ile His Ala Ile Ile Asn Ile
                260
                                    265
```

VERIFICATION SUMMARYDATE: 04/24/2003PATENT APPLICATION:US/10/043,639ATIME: 16:21:34

input Set : D:\Chep004.app

Output Set: N:\CRF4\04242003\J043639A.raw

L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date